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Range: from to Reverse complemented strand Features: Refresh

□ 1: Y08593. Reports H.sapiens mRNA for [gi:1834496]

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Comment Features Sequence

LOCUS Y08593 363 bp mRNA linear PRI 11-MAR-1999
 DEFINITION H.sapiens mRNA for Ig heavy chain anti-TTd, variable region.
 ACCESSION Y08593
 VERSION Y08593.1 GI:1834496
 KEYWORDS anti-TTd gene; immunoglobulin variable region heavy chain.
 SOURCE Homo sapiens (human)
 ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 Mammalia; Eutheria; Euarchontoglires; Primates; Haplorrhini;
 Catarrhini; Hominidae; Homo.
 REFERENCE 1
 AUTHORS Mersmann,M., Schmidt,A., Tesar,M., Schoneberg,A., Welschof,M.,
 Kipriyanov,S., Terness,P., Little,M., Pfizenmaier,K. and
 Moosmayer,D.
 TITLE Monitoring of scFv selected by phage display using detection of
 scFv-pIII fusion proteins in a microtiter scale assay
 JOURNAL J. Immunol. Methods 220 (1-2), 51-58 (1998)
 PUBMED 9839925
 REFERENCE 2 (bases 1 to 363)
 AUTHORS Welschof,M.
 TITLE Direct Submission
 JOURNAL Submitted (02-OCT-1996) M. Welschof, Universitaet Heidelberg,
 Institute of Immunology, Department of Transplantation Immunology,
 INF 305, 69120 Heidelberg, FRG
 COMMENT Related sequence: Z14202.
 FEATURES Location/Qualifiers
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 SPYWYLDLWGRGTLVTVSS"

ORIGIN
1 gaggtgcagc tggtgagtc tgggggaggc ttgggacagc cgggggggtc cctgagactc
61 tcctgtgcag ccaccggatt cagttcaac aactatgcca tgagttggc ccgtcaggct
121 ccagggaaagg ggctggagtg ggtctcgagt attagtggtg gtgggtggtag tagatactat
181 gcagactccq tqaaggqccq qttcaccatc tccaaqaqaca qttccaaqaaa cacqgtgtat

241 ctgcaaatga acaggctgag agccgaggac acggccgcat attactgtgc gagatacagt
301 agtggctccc cttattggta tctcgatctg tggggccgtg gcaccctggt cactgtctcc
361 tca

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241 cctgaagatt ttgcagtgt a ttactgtcag cagtatggta gctcaccccg aactttcggc
301 cctgggacca aagtggagat caaa

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